

## Exhibit 1

Blast Result

Page 1 of 1

 NCBI Blast 2 Sequences results

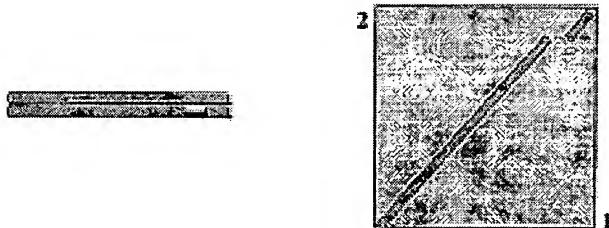
PubMed Entrez BLAST OMIM Taxonomy Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.17 [Aug-19-2007]

Matrix BLOSUM62  gap open: 11  gap extension: 1   
X-dropoff: 0  expect: 10.000 wordsize: 3  Filter  View option Standard   
Masking character option: X for protein, n for nucleotide  Masking color option: Black   
 Show CDS translation

Sequence 1: unnamed protein product  
Length = 71 (1 .. 71)

Sequence 2: unnamed protein product  
Length = 64 (1 .. 64)



NOTE: Bitscore and expect value are calculated based on the size of the nr database.

[REDACTED]

Score = 94.0 bits (232), Expect = 2e-18  
Identifications = 60/70 (85%), Positives = 61/70 (87%), Gaps = 2/70 (10%)

Query 1:	MVLRQLSRKAVSVKVKYETNGTAKRACGILILFLLDFLINFCTGEGKQQRQESLTDQ	60
	MVLRQLSRKAVSVKVKYETNGTAKRACGILILFLLDFLINFCTGEGKQQRQESLTDQ	60
Subject 1:	MVLRQLSRKAVSVKVKYETNGTAKRACGILILFLLDFLINFCTGEGKQQRQESLTDQ	56
Query 61:	TYBALPEPKA 70 SEQ. IDN# /	
	ALPEPKA	
Subject 57:	---ALPEPKA 63 SEQ. IDN# /3	

CPU time: 0.03 user secs., 0.01 sys. secs. 0.04 total secs.